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Substitute for form 1449A/PTO (modified) INFORMATION DISCLOSURE STATEMENT BY APPLICANT (use as many sheets as necessary)				Complete if known	
Application Number		Filed concurrently herewith		J1046 U.S. PTO 09/28/01	
Filing Date		September 28, 2001			
First Named Inventor		Eric C. Hannah			
Group Art Unit		Not yet assigned			
Examiner Name		Not yet assigned			
Attorney Docket Number		42390P11816			
Sheet	1	of	2		

U.S. PATENT DOCUMENTS						
Examiner Initials	Cite No. ¹	U.S. Patent Document Number	Kind Code ² (if known)	Name of Patentee or Applicant of Cited Document	Date of Publication of Cited Document MM-DD-YYYY	Pages, Columns, Lines Where Relevant Passages or Figures Appear
MB		5,436,850		Eisenberg et al.	07-25-1995	
		5,600,571		Friesner et al.	02-04-1997	
		5,680,319		Rose et al.	10-21-1997	
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MB		5,884,230		Srinivasan et al.	03-16-1999	

FOREIGN PATENT DOCUMENTS								
Examiner Initials*	Cite No. ¹	Office ³	Foreign Patent Number ⁴	Kind Code ⁵ (if known)	Name of Patentee or Applicant of Cited Document	Date of Publication of Cited Document MM-DD-YYYY	Pages, Columns, Lines Where Relevant Passages or Relevant Figures Appear	T ⁶
			WO 98/48270		California Institute of Technology	10-29-1998		
Examiner Signature					/Michael Borin/		Date Considered	06/07/2006

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 J1046 U.S. PTO
 09/26/02

10/02/00

OTHER PRIOR ART - NON PATENT LITERATURE DOCUMENTS

Examiner Initials	Cite No. ¹	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date page(s), volume-issue number(s), publisher, city and/or country where published.	T ²
MB		Al-Lazikani, Bissan <i>et al.</i> , "Protein structure prediction", <i>Curr. Opin. in Chem. Biol.</i> , 5:51-56, 2001	
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		Debe, Derek A. <i>et al.</i> , "The topomer-sampling model of protein folding", <i>Proc. Natl. Acad. Sci USA</i> , 96:2596-2601, March 1999	
		Debe and Goddard III, "First Principles Prediction of Protein Folding Rates", <i>J. Mol. Biol.</i> , 294:619-625, 1999	
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		Pillardy, Jaroslaw <i>et al.</i> , "Recent improvements in prediction of protein structure by global optimization of a potential energy function", <i>PNAS</i> 98(5) 2329-2333, February 27, 2001	
MB		Thomas and Dill, "An iterative method for extracting energy-like quantities from protein structures", <i>Proc. Natl. Acad. Sci USA</i> 93:11628-11633, October 1996	

Examiner Signature	/Michael Borin/	Date Considered	06/07/2005
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